



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/888,233
Source: OIPE
Date Processed by STIC: 7/2/2001

BEST AVAILABLE COPY

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25. Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/888,233

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- Numbering
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0 A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- "bug"
- 7 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (OLD RULES) (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 (NEW RULES) <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
 (NEW RULES) Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213> Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or
 Response scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (Sec "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0 Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
 "bug" resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

OIKE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/888,233

DATE: 07/07/2001

TIME: 12:52:58

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Does Not Comply
Corrected Diskette Needed

P.4

3 <110> APPLICANT: BLAKELY, RANDY D.
 4 NASS, RICHARD
 5 MILLER, DAVID M.
 7 <120> TITLE OF INVENTION: ASSAY FOR TOXIN INDUCED NEURONAL DEGENERATION AND
 8 VIABILITY IN C. ELEGANS
 10 <130> FILE REFERENCE: VBLT:007US
 11 <140> CURRENT APPLICATION NUMBER: UNKNOWN
 12 <141> CURRENT FILING DATE: 2001-06-22
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 16 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

P.4

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208 caacaatagt cagacctgt ctctacaaga aaacaacaac aacaacaaa aaacacacac 9900
209 aaaaataact tagccggtgc tgtggtgcac acctgtagtc tcagctgctc aggaggctga 9960
210 ggtgcaagg tcaactgaac ctaggaggtt gaggcagtga gttgtaatca caactgtatt 10020
211 ccatcctggg tgacagagcg agacctcatg ttaaaaaaga aaaaaaaaag aaaaaagaat 10080
212 acagatgaac agtcatgaag acattattga atgctcttag aagattgtaa aattgctctc 10140
213 tgggaagtgtg ggggaagggtg gaagtgatat ccatgcattg ttagtagaaa gccacgctga 10200
214 agctcacaca gccttgcaact ttgataggag tggggagggg tgcaggggaa ggaggagcaa 10260
215 accagagtgt ctgtcttgag gcctccatgg gccagtgcc cagccctgtg gtgagggtg 10320
216 gcattcccag ctcccggtgc ccagctgtac catctccagg cgtgagaagc acccatcctt 10380
217 tcccagagga atgcccgtga atgcttcggg gtctgccatc cgcaacaggt atgtccctag 10440
218 ccctggctga tgaattgttg cgttctctgt gtgtgtttat ttttcatatt ggctgaagac 10500

```

see item 9
on Enon
summary
sheet

RAW SEQUENCE LISTING

DATE: 07/07/2001

PATENT APPLICATION: US/09/888,233

TIME: 12:52:58

Input Set : A:\Vblt007.app

Output Set: N:\CRF3\07062001\I888233.raw

```
219 caagagggaa gaagcacaga attctcaact cccagtgtgc ccatgagtaa gagcaaatgc 10560
220 tccgtgggac tcatgtcttc cgtggtggcc ccggctaagg agcccaatgc cgtgggcccg 10620
221 aaggaggtgg agctcatcct tgtcaaggag cagaacggag tgcagttcac cagttccacc 10680
222 ttcaccaacc cgcggcagag ccccgaggag gcccaggatc gggagacctg gggcaagaag 10740
223 atcgatttct cctgtccgtc attggctttg ctgtggacct ggccaacgtc tggcggttcc 10800
224 cctacctgtg ctacaaaaat ggtggcggta atcccatctc agcttccctg a 10851
```

VERIFICATION SUMMARY

DATE: 07/07/2001

PATENT APPLICATION: US/09/888,233

TIME: 12:52:59

Input Set : A:\Vblt007.app

Output Set: N:\CRF3\07062001\I888233.raw

L:11 M:283 W: Missing Blank Line separator, <140> field identifier

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number

L:202 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2

M:340 Repeated in SeqNo=2